## **FORM PTO-1449**

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U.S. PATENT DOCUMENTS Ref. Document Subaminer Filing Date Initial No Number Date Name Class Class FOREIGN PATENT DOCUMENTS PCT-98/50555 A3 11/1998 -AA CAL OTHER DOCUMENTS (Including Author, Title, Date, etc.) A. van Belkum et al., "Outbreak of Amoxicillin-Resistant Haemophilus influenza Type AB b. Variable Number of Tandem Repeats as Novel Molecular Markers," Journal of cor Clinical Microbiology, 35(6):1517-1520 (1997). A. van Belkum et al., "Variable Number of Tandem Repeats in Clinical Strains of AC Haemophilus influenza," Infection and Immunity, 65(12):5017-5027 (1997). A. van Belkum et al., "Short-Sequence DNA Repeats in Prokaryotic Genomes," AD Microbiology and Molecular Biology Reviews, 62(2):275-293 (1998). AE N. Kobayashi et al., "Analysis of genomic diversity within the Xr-region of the protein A gene in clinical isolates of Staphylococcus aureus," Epidemiol. Infect., 122:241-249 (1999).Database EMBL 'Online!, AAX13101 "Enterococcus faecalis genome contig SEQ ID AF NO:164, March 19, 1999. B. Shopsin et al., "Evaluation of Protein A Gene Polymorphic Region DNA Sequencing AG for Typing of Staphylococcus aureus Strains," J. Clin. Micr., 3556-3563 (1999). AH B. Shopsin et al., "Use of Coagulase Gene (coa) Repeat Region Nucleotide Sequences for Typing of Methicillin-Resistant Staphylococcus aureus Strains," J. Clin. Microbiol., 38(9):3453-3456 (2000). ΑI A. van Belkum, "Short sequence repeats in microbial pathogenesis and evolution," Cell. Mol. Life Sci., 56:729-734 (1999). A. van Belkum, "The role of short sequence repeats in epidemiologic typing," Current ΑJ Opinion in Microbiol., 2(3) 306-311 (1999).

Examiner	Uller	Date Considered	3/30/64	
Examiner: Initial if reference consider, whether or not citation is in conformance with MPEP §609.  Draw line through citation if not in conformance and not considered.				
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